

§Appl. No. 10/030,688
Amdt. dated November 24, 2004
Reply to Office Action of, August 24, 2004

In the Specification:

Please amend the specification as follows:

Other programs for determining identity and/or similarity between sequences are also known in the art, for instance the BLAST family of programs (Altschul S F et al. J Mol Biol, 215, 403-410, 1990 Altschul S F et al, Nucleic Acids res., 25:389-3402, 1997, available from the National Center for Biotechnology Information (NCBI), Bethesda, Maryland, USA and accessible through the home page of the NCBI at ~~www.ncbi.nih.gov~~ the world wide web at ncbi.nlm.nih.gov) and FASTA (Pearson W R, Methods in Enzymology, 183, 63-99, 1990; Pearson W R and Lipman D J, Proc Nat Acad Sci USA, 85, 2444-2448, 1988, available as part of the Wisconsin Sequence Analysis Package).